

FIG. 1

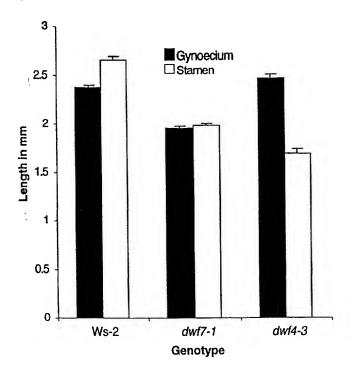


FIG. 2

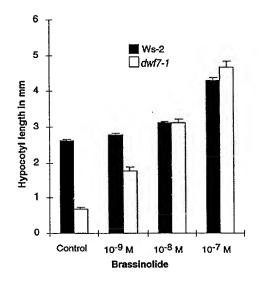


FIG. 3

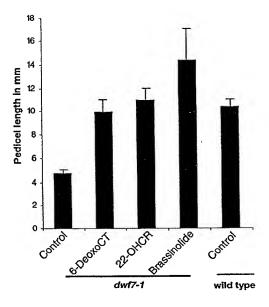
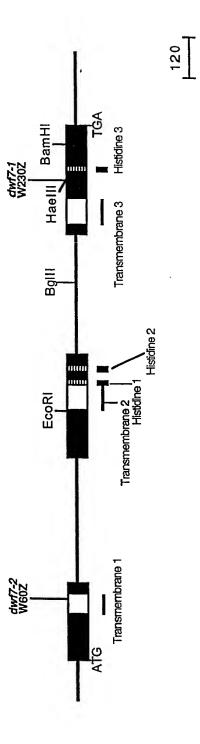


FIG. 4

F/G. 5



F/G. 6

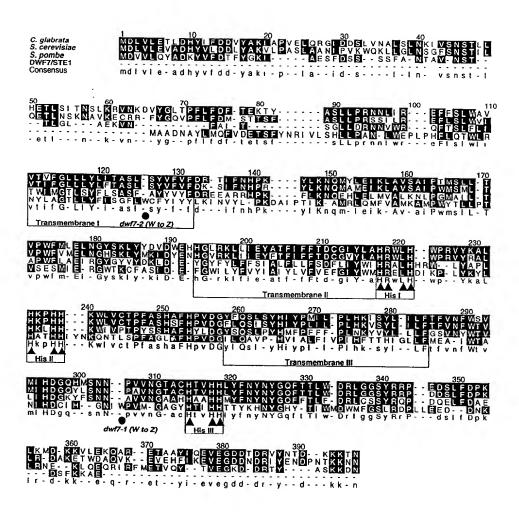


FIG. 7

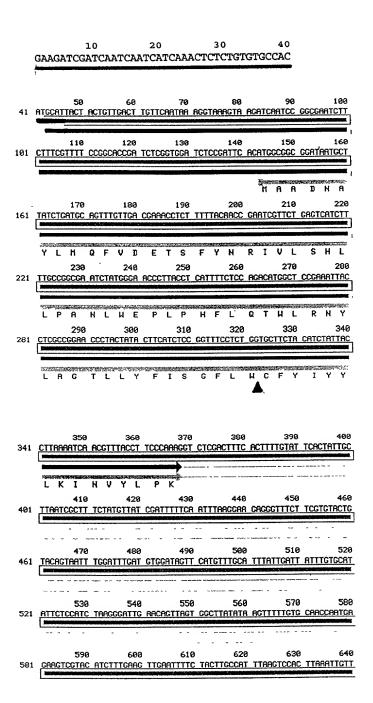


FIG. 8A

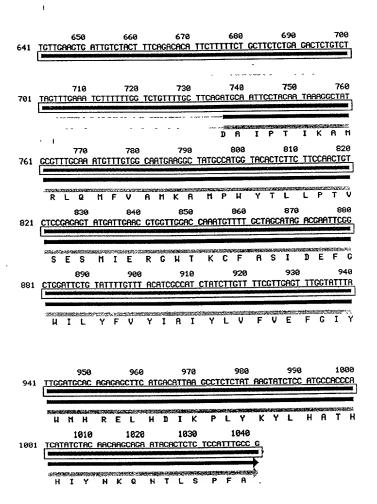


FIG. 8B

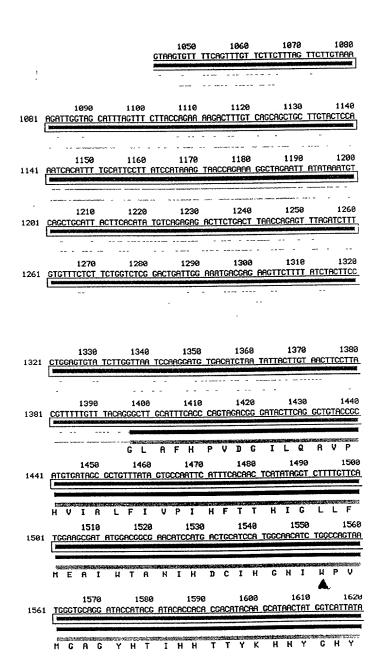


FIG. 8C

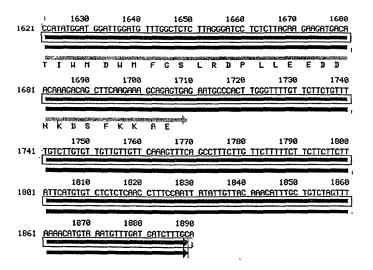


FIG. 8D

1	MAADNAYLMQ	FVDETSFYNR	IVLSHLLPAN	LWEPLPHFLQ	TWLRNYLAG
51	LLYFISGFLW	CFYIYYLKIN	VYLPKDAIPT	IKAMRLQMFV	AMKAMPWYTI
101	LPTVSESMIE	RGWTKCFASI	DEFGWILYFV	YIAIYLVFVE	FGIYWMHREI
151	HDIKPLYKYL	HATHHIYNKQ	NTLSPFAGLA	FHPVDGILQA	VPHVIALFI
201	PIHFTTHIGL	LFMEAIWTAN	IHDCIHGNIW	PVMGAGYHTI	ннттукниус
251	HYTIWMDWMF	GSLRDPLLEE	DDNKDSFKKA	E	

FIG. 9

10	30	50									
GTTTGGTATTTATTGGATGCCAAACCATAAATAACCTACC	CACAGAGAGCTTCATGACA GTGTCTCTCGAAGTACTGT	TTAAGCCTCTCTATAAGTATCT AATTCGGAGAGATATTCATAGA									
70	90	110									
CCATGCCACCCATCATATC	TACAACAAGCAGAATACAC ATGTTGTTCGTCTTATGTG	TCTCTCCATTTGCCGGTAAGTG AGAGAGGTAAACGGCCATTCAC									
130	150	170									
TTTTCAGTTTGTTCTTCTT AAAAGTCAAACAAGAAGAA	TAGTTCTTGTAAAAGATTG ATCAAGAACATTTTCTAAC	GTAGCATTTAGTTTCTTACCAG CATCGTAAATCAAAGAATGGTC									
190	210	230									
AAAAGACTTTGTCAGCAGCTTTTTCTGAAACAGTCGTCGA	IGCTTGTACTCCAAATCAC. ACGAACATGAGGTTTAGTG	ATTTTGCATTCCTTATCCATAA TAAAACGTAAGGAATAGGTATT									
250	270	290									
310	330	350									
AGACTTCTGACTTAACCAGA TCTGAAGACTGAATTGGTCT	AGACTTCTGACTTAACCAGAGTTTAGATCTTTGTGTTTCTCTTCTGGTCTCGGACTGATT TCTGAAGACTGAATTGGTCTCAAATCTAGAAACACAAAGAGAAGACCAGAGCCTGACTAA										
370	390	410									
GGAAATGACGAGAAGTTCTT	FTTATCTACTTCCCTGGAG: AAATAGATGAAGGGACCTC	IGTATCTTGGTTAATCCAAGGA ACATAGAACCAATTAGGTTCCT									
430	450	470									
TGTGACATCTAAATATTACT ACACTGTAGATTTATAATGA	PTGTAACTTCCTTACGTTT AACATTGAAGGAATGCAAA!	TTGTTTACAGGGCTTGCATTCA AACAAATGTCCCGAACGTAAGT									
490	510	530									
CCCAGTAGACGGGATACTTA GGGTCATCTGCCCTATGAAT	AAGGCTGTACCGCATGTGAT TTCCGACATGGCGTACACTA	FAGCGCTGTTATAGTGCCAATT ATCGCGACAATATCACGGTTAA									
550	570	590									
CATTTCACAACTCATATAGG GTAAAGTGTTGAGTATATCC	GTCTTTTGTTCATGGAAGCC	GATATGGACGGCGAACATCCAT									

FIG. 10A

610	630	650
		AGGATACCATACGATACACCAC ICCTATGGTATGCTATGTGGTG
670	690	710
		GATGGATTGGATGTTTGGCTCT CTACCTAACCTACAAACCGAGA
730	750	770
		CAGCTTCAAGAAAGCAGAGTGA GTCGAAGTTCTTTCGTCTCACT
790	810	830
		TGTTGTTGTTGTTCAAAGTTTC ACAACAACAACAAGTTTCAAAG
850	870	890
		TGTCTCTCTCAACCTTTCCAAT ACAGAGAGAGTTGGAAAGGTTA
910	930	950
		GTAAATGTTTGATGATCTTTGC CATTTACAAACTACTAGAAACG
970	990	1010
		TAGATTGTCGATTGTTGGTATT ATCTAACAGCTAACAACCATAA
1030	1050	1070
		TGACCAGTCCGGCTTAACCACC ACTGGTCAGGCCGAATTGGTGG
1090	1110	1130
		GCCCCAATATATAGATGGGCCA CGGGGTTATATATCTACCCGGT
1150	1170	1190
		ACAGTTAGACACCTGCTAATTA TGTCAATCTGTGGACGATTAAT

1210		1230		1250									
1270		1290		1310									
TAGTAAACGCAATTTAACCCTTATAAGTTTAATCGTATTCAACGAATGACCCAGAGACT ATCATTTGCGTTAAATTGGGAATATTCAAATTAGCATAAGTTGCTTACTGGGTCTCTGA													
1330		1350		1370	1370								
TAAATAAATCCATCGTAACCCTCCACTTCAAAATTCTTTTTAAAAAAGTAGCAAATCATTTATTT													
1390		1410		1430									
1450		1470		1490	1490								
1510		1530		1550									
GTTTCATGGCGGCCCCCCCCCCCCCCCCCCCCCCCCCCC			TAGTCTAGCAG										
1570		1590		1610									
ACAACCGAAtGGTT TGTTGGCTTaCCA/ N R M V		_	ACTTAGATACC										
1630		1650		1670									
TCCTCCAGACATGC AGGAGGTCTGTACC L Q T W		GAGCGGC L A G	CTTTGTATGAG	ATGAAGTAG Y F I									
1690		1710		1730	•								
TCCTCTGGTGCTTC		GGAATTTG	AGTTGCAAATG										

		175	0					:	L770	)			1	1790										
						-													TATC ATAG					
		181	.0					:	1830	0					18	350								
																			IGTC ACAG					
-		CTATCTAACACAATATGCAATTGGAAAAAAAGAAT 1870 1890													1910									
		193	30										1	970										
		1990 2010										2030												
																			GAGA CTCT R					
		20!	50						207	0					2	090								
																			TCTT AGAA L					
		21	10						213	0					2	150								
																			TGAC ACTG D					
		21	70						219	0					2	210								
	AAA	GTT	GAC	CAA	.GGA		AAA	GGA	GAT	GTA	TCG	AGA	GAT.	ACA	ACA	AAA	TCA	ACT	GTTt CAAa F					
		22	30						225	0					2	270								
	CTA	AAT	AAC	CCA	AGT		TCT	CGA	AGT	'ACT	'GTA	ATT	TAA						CCAT .GGTA					

		2290								.0				:	2330						
													CTC( GAG(								
A	T	H	H	М	Y	N	K	Q	N	T	L	S	P	F	A	JOC,	⊖F1±1	nc <sub>E</sub>	011	10	
		23	50						237	70						239	0				
													ATGO								
	2410 2430 2450																				
														GC							
AAA	AAG	CCC	'AAZ	\AG'l	CCC G	CGA( L	GCG.	TAA( F	3GT∤ H	AGG( P	CGA: L	CCT D	GCC G	CTA' I	TGA L	AGT Q	CCG. A	ATA I	TGG P	CG H	
		24	70						249	90						251	0				
	CAC												AAC TTG T								
		25	30						25!	50						257	0				
ACC	CTT	CCC	TAT	raco	CTG	TCG'	TTC	GTA	GGT	ACT.	AAC	GTA	ACA TGT	ACC	aTT	GTA	GAC	CGG	ATA		
Ε	3	G	Ι	W	Т	A	S	Ι	Н	D	С	I	Η	G	N	Ι	M	Ρ	Ι	M	
		25	90						26	10						263	0				
													.CAA 'GTT								
		A	G	Y	H	T	I	Н	H	T	T	Y	K	Н	N	Y	G	H	Υ	Т	
		26	550						26	70						269	0				
													TCC								
	IA1	W W	M M	D	JAC W	CTA M	CAA F	ACC G	GAG. S	AGA L	A'I'A <b>M</b>	V.	AGG P	AAA L	A.	E ETCT	TTT K	D TCT	GTC S	'AA F	
		27	710						27	30						275	0				
AG*													CATG STAC								
		27	770						27	90						281	.0				
													GTG CAC			-					

FIG. 10E

 ${\tt ATTATATATTGcTGGATGAAGAGTTCAAATTTGGACTAAATCTG}\\ {\tt TAATATATTAACgACCTACTTCTCAAGTTTAAACCTGATTTAGAC}\\$ 

maatmadynd qivnetsfyn rmvlshllpv nlweplphfl qtwlrnylag nilyfisgfl wcfyiyylkl nvyvpkesip trkamllqiy vamkampwyt llpavseymi ehgwtkcyst ldhfnwflcf lyialylvlv efmiywvhke lhdikflykh lhathhmynk qntlspfagl afhpldgilq aiphvialfi vpihlithls llflegiwta sihdcihgni wpimgagyht ihhttykhny ghytiwmdwm fgslmvplae kdsfkekek